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OIPE

## RAW SEQUENCE LISTING

DATE: 01/28/2002

PATENT APPLICATION: US/10/010,229

TIME: 11:32:05

Input Set : N:\Crif3\RULE60\10010229.raw

Output Set: N:\CRF3\01282002\J010229.raw

1 <110> APPLICANT: Le, Junming  
 2 Vilcek, Jan  
 3 Daddona, Peter  
 4 Ghrayeb, John  
 5 Knight, David M.  
 6 Siegel, Scott  
 7 <120> TITLE OF INVENTION: Anti-TNF Antibodies and Peptides of  
 8 Human Tumor Necrosis Factor  
 9 <130> FILE REFERENCE: 0975.1005-013  
 10 <140> CURRENT APPLICATION NUMBER: 10/010,229  
 11 <141> CURRENT FILING DATE: 2001-12-07  
 12 <150> PRIOR APPLICATION NUMBER: US/09/927,703  
 13 <151> PRIOR FILING DATE: 2001-08-10  
 15 <160> NUMBER OF SEQ ID NOS: 19  
 16 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 18 <210> SEQ ID NO: 1  
 19 <211> LENGTH: 157  
 20 <212> TYPE: PRT  
 21 <213> ORGANISM: Homo sapiens  
 22 <400> SEQUENCE: 1  
 23 Val Arg Ser Ser Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val  
 24 1 5 10 15  
 25 Val Ala Asn Pro Gln Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg  
 26 20 25 30  
 27 Ala Asn Ala Leu Leu Ala Asn Gly Val Glu Leu Arg Asp Asn Gln Leu  
 28 35 40 45  
 29 Val Val Pro Ser Glu Gly Leu Tyr Leu Ile Tyr Ser Gln Val Leu Phe  
 30 50 55 60  
 31 Lys Gly Gln Gly Cys Pro Ser Thr His Val Leu Leu Thr His Thr Ile  
 32 65 70 75 80  
 33 Ser Arg Ile Ala Val Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala  
 34 85 90 95  
 35 Ile Lys Ser Pro Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys  
 36 100 105 110  
 37 Pro Trp Tyr Glu Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys  
 38 115 120 125  
 39 Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg Pro Asp Tyr Leu Asp Phe  
 40 130 135 140  
 41 Ala Glu Ser Gly Gln Val Tyr Phe Gly Ile Ile Ala Leu  
 42 145 150 155  
 44 <210> SEQ ID NO: 2  
 45 <211> LENGTH: 321  
 46 <212> TYPE: DNA

ENTERED

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47 <213> ORGANISM: Mus Balb/c
48 <220> FEATURE:
49 <221> NAME/KEY: CDS
50 <222> LOCATION: (1)...(321)
51 <400> SEQUENCE: 2
52     gac atc ttg ctg act cag tct cca gcc atc ctg tct gtg agt cca gga      48
53     Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
54     1           5           10           15
55     gaa aga gtc agt ttc tcc tgc agg gcc agt cag ttc gtt ggc tca agc      96
56     Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser
57           20           25           30
58     atc cac tgg tat cag caa aga aca aat ggt tct cca agg ctt ctc ata      144
59     Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
60           35           40           45
61     aag tat gct tct gag tct atg tct ggg atc cct tcc agg ttt agt ggc      192
62     Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly
63           50           55           60
64     agt gga tca ggg aca gat ttt act ctt agc atc aac act gtg gag tct      240
65     Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser
66           65           70           75           80
67     gaa gat att gca gat tat tac tgt caa caa agt cat agc tgg cca ttc      288
68     Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
69           85           90           95
70     acg ttc ggc tcg ggg aca aat ttg gaa gta aaa      321
71     Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys
72           100           105
74 <210> SEQ ID NO: 3
75 <211> LENGTH: 107
76 <212> TYPE: PRT
77 <213> ORGANISM: Mus Balb/c
78 <400> SEQUENCE: 3
79     Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
80     1           5           10           15
81     Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser
82           20           25           30
83     Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
84           35           40           45
85     Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly
86           50           55           60
87     Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser
88           65           70           75           80
89     Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
90           85           90           95
91     Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys
92           100           105
94 <210> SEQ ID NO: 4
95 <211> LENGTH: 357
96 <212> TYPE: DNA
97 <213> ORGANISM: Mus Balb/c

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98 &lt;220&gt; FEATURE:

99 &lt;221&gt; NAME/KEY: CDS

100 &lt;222&gt; LOCATION: (1)...(357)

101 &lt;400&gt; SEQUENCE: 4

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102      gaa gtg aag ctt gag gag tct gga gga ggc ttg gtg caa cct gga gga      48
103      Glu Val Lys Leu Glu Glu Ser Gly Gly Leu Val Gln Pro Gly Gly
104      1          5          10          15
105      tcc atg aaa ctc tcc tgt gtt gcc tct gga ttc att ttc agt aac cac      96
106      Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His
107      20          25          30
108      tgg atg aac tgg gtc cgc cag tct cca gag aag ggg ctt gag tgg gtt      144
109      Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val
110      35          40          45
111      gct gaa att aga tca aaa tct att aat tct gca aca cat tat gcg gag      192
112      Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu
113      50          55          60
114      tct gtg aaa ggg agg ttc acc atc tca aga gat gat tcc aaa agt gct      240
115      Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala
116      65          70          75          80
117      gtc tac ctg caa atg acc gac tta aga act gaa gac act ggc gtt tat      288
118      Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr
119      85          90          95
120      tac tgt tcc agg aat tac tac ggt agt acc tac gac tac tgg ggc caa      336
121      Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln
122      100          105          110
123      ggc acc act ctc aca gtc tcc      357
124      Gly Thr Thr Leu Thr Val Ser
125      115

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127 &lt;210&gt; SEQ ID NO: 5

128 &lt;211&gt; LENGTH: 119

129 &lt;212&gt; TYPE: PRT

130 &lt;213&gt; ORGANISM: Mus Balb/c

131 &lt;400&gt; SEQUENCE: 5

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132      Glu Val Lys Leu Glu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
133      1          5          10          15
134      Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His
135      20          25          30
136      Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val
137      35          40          45
138      Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu
139      50          55          60
140      Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala
141      65          70          75          80
142      Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr
143      85          90          95
144      Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln
145      100          105          110
146      Gly Thr Thr Leu Thr Val Ser
147      115

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Input Set : N:\CrF3\RULE60\10010229.raw  
 Output Set: N:\CRF3\01282002\J010229.raw

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149 <210> SEQ ID NO: 6
150 <211> LENGTH: 8
151 <212> TYPE: PRT
152 <213> ORGANISM: Homo sapiens
153 <400> SEQUENCE: 6
154     Gly Thr Leu Val Thr Val Ser Ser
155         1             5
157 <210> SEQ ID NO: 7
158 <211> LENGTH: 7
159 <212> TYPE: PRT
160 <213> ORGANISM: Homo sapiens
161 <400> SEQUENCE: 7
162     Gly Thr Lys Leu Glu Ile Lys
163         1             5
165 <210> SEQ ID NO: 8
166 <211> LENGTH: 20
167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: PCR oligonucleotides
171 <400> SEQUENCE: 8
172     cctggatacc tgtgaaaaga
174 <210> SEQ ID NO: 9
175 <211> LENGTH: 27
176 <212> TYPE: DNA
177 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: PCR oligonucleotides
180 <400> SEQUENCE: 9
181     cctggtacct tagtcaccgt ctctca
183 <210> SEQ ID NO: 10
184 <211> LENGTH: 27
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
188 <223> OTHER INFORMATION: PCR oligonucleotides
189 <400> SEQUENCE: 10
190     aatagatatc tccttcaaca cctgcaa
192 <210> SEQ ID NO: 11
193 <211> LENGTH: 21
194 <212> TYPE: DNA
195 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: PCR oligonucleotides
198 <400> SEQUENCE: 11
199     atcgggacaa agttggaaat a
201 <210> SEQ ID NO: 12
202 <211> LENGTH: 16
203 <212> TYPE: DNA

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Input Set : N:\Crf3\RULE60\10010229.raw

Output Set: N:\CRF3\01282002\J010229.raw

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204 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: PCR oligonucleotides
207 <400> SEQUENCE: 12
208      ggcggtcttg taccgg                                16
210 <210> SEQ ID NO: 13
211 <211> LENGTH: 19
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: PCR oligonucleotides
216 <400> SEQUENCE: 13
217      gtcaacaaca tagtcatca                                19
219 <210> SEQ ID NO: 14
220 <211> LENGTH: 23
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: PCR oligonucleotides
225 <400> SEQUENCE: 14
226      cacaggtgtg tccccaagga aaa                            23
228 <210> SEQ ID NO: 15
229 <211> LENGTH: 18
230 <212> TYPE: DNA
231 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: PCR oligonucleotides
234 <400> SEQUENCE: 15
235      aatctggggg aggcacaa                                18
237 <210> SEQ ID NO: 16
238 <211> LENGTH: 17
239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: PCR oligonucleotides
243 <400> SEQUENCE: 16
244      agtgtgtgtc cccaagg                                17
246 <210> SEQ ID NO: 17
247 <211> LENGTH: 24
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: PCR oligonucleotides
252 <400> SEQUENCE: 17
253      cacagctgcc cgcccaggtg gcat                            24
255 <210> SEQ ID NO: 18
256 <211> LENGTH: 17
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/010,229

DATE: 01/28/2002

TIME: 11:32:06

Input Set : N:\Crf3\RULE60\10010229.raw

Output Set: N:\CRF3\01282002\J010229.raw